RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10 518,184

Source: PCT

Date Processed by STIC: 12-28-04

ENTERED

PCT

RAW SEQUENCE LISTING   DATE: 12/28/2004
PATENT APPLICATION: US/10/518,184   TIME: 11:53:52

Input Set : A:\3057USOP.SEQ.txt
Output Set: N:\CRF4\12282004\J518184.raw

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     3  <110> APPLICANT: WATANABE, TATSUYA
     4        INAZUKA, MASAKAZU
     6  <120> TITLE OF INVENTION: Prophylactic/therapeutic agents for bone or joint diseases
     8  <130> FILE REFERENCE: 3057 USOP
C--> 10  <140> CURRENT APPLICATION NUMBER: US/10/518,184
C--> 11  <141> CURRENT FILING DATE: 2004-12-16
    13  <150> PRIOR APPLICATION NUMBER: PCT/JP03/007741
W--> 14  <151> PRIOR FILING DATE: 2003-6-18
    16  <150> PRIOR APPLICATION NUMBER: JP2002-178715
W--> 17  <151> PRIOR FILING DATE: 2002-6-19
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    22  <211> LENGTH: 871
    23  <212> TYPE: PRT
    24  <213> ORGANISM: Homo sapiens
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    29  Leu Leu Gly Ser Leu Cys Ser Thr Val Arg Ser Pro Arg Phe Arg Gly
    30              20                  25                  30
    31  Arg Ile Gln Gln Glu Arg Lys Asn Ile Arg Pro Asn Ile Ile Leu Val
    32          35                  40                  45
    33  Leu Thr Asp Asp Gln Asp Val Glu Leu Gly Ser Leu Gln Val Met Asn
    34      50                  55                  60
    35  Lys Thr Arg Lys Ile Met Glu His Gly Gly Ala Thr Phe Ile Asn Ala
    36  65                  70                  75                  80
    37  Phe Val Thr Thr Pro Met Cys Cys Pro Ser Arg Ser Ser Met Leu Thr
    38              85                  90                  95
    39  Gly Lys Tyr Val His Asn His Asn Val Tyr Thr Asn Asn Glu Asn Cys
    40              100                 105                 110
    41  Ser Ser Pro Ser Trp Gln Ala Met His Glu Pro Arg Thr Phe Ala Val
    42          115                 120                 125
    43  Tyr Leu Asn Asn Thr Gly Tyr Arg Thr Ala Phe Phe Gly Lys Tyr Leu
    44      130                 135                 140
    45  Asn Glu Tyr Asn Gly Ser Tyr Ile Pro Pro Gly Trp Arg Glu Trp Leu
    46  145                 150                 155                 160
    47  Gly Leu Ile Lys Asn Ser Arg Phe Tyr Asn Tyr Thr Val Cys Arg Asn
    48              165                 170                 175
    49  Gly Ile Lys Glu Lys His Gly Phe Asp Tyr Ala Lys Asp Tyr Phe Thr
    50              180                 185                 190
    51  Asp Leu Ile Thr Asn Glu Ser Ile Asn Tyr Phe Lys Met Ser Lys Arg
    52          195                 200                 205
    53  Met Tyr Pro His Arg Pro Val Met Met Val Ile Ser His Ala Ala Pro
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55 His Gly Pro Glu Asp Ser Ala Pro Gln Phe Ser Lys Leu Tyr Pro Asn
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57 Ala Ser Gln His Ile Thr Pro Ser Tyr Asn Tyr Ala Pro Asn Met Asp
58                 245                     250                     255
59 Lys His Trp Ile Met Gln Tyr Thr Gly Pro Met Leu Pro Ile His Met
60             260                     265                     270
61 Glu Phe Thr Asn Ile Leu Gln Arg Lys Arg Leu Gln Thr Leu Met Ser
62         275                     280                     285
63 Val Asp Asp Ser Val Glu Arg Leu Tyr Asn Met Leu Val Glu Thr Gly
64     290                     295                     300
65 Glu Leu Glu Asn Thr Tyr Ile Ile Tyr Thr Ala Asp His Gly Tyr His
66 305                     310                     315                     320
67 Ile Gly Gln Phe Gly Leu Val Lys Gly Lys Ser Met Pro Tyr Asp Phe
68                 325                     330                     335
69 Asp Ile Arg Val Pro Phe Phe Ile Arg Gly Pro Ser Val Glu Pro Gly
70             340                     345                     350
71 Ser Ile Val Pro Gln Ile Val Leu Asn Ile Asp Leu Ala Pro Thr Ile
72         355                     360                     365
73 Leu Asp Ile Ala Gly Leu Asp Thr Pro Pro Asp Val Asp Gly Lys Ser
74     370                     375                     380
75 Val Leu Lys Leu Leu Asp Pro Glu Lys Pro Gly Asn Arg Phe Arg Thr
76 385                     390                     395                     400
77 Asn Lys Lys Ala Lys Ile Trp Arg Asp Thr Phe Leu Val Glu Arg Gly
78             405                     410                     415
79 Lys Phe Leu Arg Lys Lys Glu Glu Ser Ser Lys Asn Ile Gln Gln Ser
80         420                     425                     430
81 Asn His Leu Pro Lys Tyr Glu Arg Val Lys Glu Leu Cys Gln Gln Ala
82         435                     440                     445
83 Arg Tyr Gln Thr Ala Cys Glu Gln Pro Gly Gln Lys Trp Gln Cys Ile
84     450                     455                     460
85 Glu Asp Thr Ser Gly Lys Leu Arg Ile His Lys Cys Lys Gly Pro Ser
86 465                     470                     475                     480
87 Asp Leu Leu Thr Val Arg Gln Ser Thr Arg Asn Leu Tyr Ala Arg Gly
88             485                     490                     495
89 Phe His Asp Lys Asp Lys Glu Cys Ser Cys Arg Glu Ser Gly Tyr Arg
90         500                     505                     510
91 Ala Ser Arg Ser Gln Arg Lys Ser Gln Arg Gln Phe Leu Arg Asn Gln
92     515                     520                     525
93 Gly Thr Pro Lys Tyr Lys Pro Arg Phe Val His Thr Arg Gln Thr Arg
94     530                     535                     540
95 Ser Leu Ser Val Glu Phe Glu Gly Glu Ile Tyr Asp Ile Asn Leu Glu
96 545                     550                     555                     560
97 Glu Glu Glu Glu Leu Gln Val Leu Gln Pro Arg Asn Ile Ala Lys Arg
98             565                     570                     575
99 His Asp Glu Gly His Lys Gly Pro Arg Asp Leu Gln Ala Ser Ser Gly
100             580                     585                     590
101 Gly Asn Arg Gly Arg Met Leu Ala Asp Ser Ser Asn Ala Val Gly Pro
102             595                     600                     605
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103 Pro Thr Thr Val Arg Val Thr His Lys Cys Phe Ile Leu Pro Asn Asp
104     610             615             620
105 Ser Ile His Cys Glu Arg Glu Leu Tyr Gln Ser Ala Arg Ala Trp Lys
106 625             630             635             640
107 Asp His Lys Ala Tyr Ile Asp Lys Glu Ile Glu Ala Leu Gln Asp Lys
108             645             650             655
109 Ile Lys Asn Leu Arg Glu Val Arg Gly His Leu Lys Arg Arg Lys Pro
110             660             665             670
111 Glu Glu Cys Ser Cys Ser Lys Gln Ser Tyr Tyr Asn Lys Glu Lys Gly
112     675             680             685
113 Val Lys Lys Gln Glu Lys Leu Lys Ser His Leu His Pro Phe Lys Glu
114     690             695             700
115 Ala Ala Gln Glu Val Asp Ser Lys Leu Gln Leu Phe Lys Glu Asn Asn
116 705             710             715             720
117 Arg Arg Arg Lys Lys Glu Arg Lys Glu Lys Arg Arg Gln Arg Lys Gly
118             725             730             735
119 Glu Glu Cys Ser Leu Pro Gly Leu Thr Cys Phe Thr His Asp Asn Asn
120     740             745             750
121 His Trp Gln Thr Ala Pro Phe Trp Asn Leu Gly Ser Phe Cys Ala Cys
122     755             760             765
123 Thr Ser Ser Asn Asn Asn Thr Tyr Trp Cys Leu Arg Thr Val Asn Glu
124     770             775             780
125 Thr His Asn Phe Leu Phe Cys Glu Phe Ala Thr Gly Phe Leu Glu Tyr
126 785             790             795             800
127 Phe Asp Met Asn Thr Asp Pro Tyr Gln Leu Thr Asn Thr Val His Thr
128             805             810             815
129 Val Glu Arg Gly Ile Leu Asn Gln Leu His Val Gln Leu Met Glu Leu
130             820             825             830
131 Arg Ser Cys Gln Gly Tyr Lys Gln Cys Asn Pro Arg Pro Lys Asn Leu
132     835             840             845
133 Asp Val Gly Asn Lys Asp Gly Gly Ser Tyr Asp Leu His Arg Gly Gln
134     850             855             860  .
135 Leu Trp Asp Gly Trp Glu Gly
136 865             870
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139 <211> LENGTH: 2613
140 <212> TYPE: DNA
141 <213> ORGANISM: Homo sapiens
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146 atccgaccca acattattct tgtgcttacc gatgatcaag atgtggagct ggggtccctg    180
147 caagtcatga acaaaacgag aaagattatg gaacatgggg gggccacctt catcaatgcc    240
148 tttgtgacta cacccatgtg ctgcccgtca cggtcctcca tgctcaccgg gaagtatgtg    300
149 cacaatcaca atgtctacac caacaacgag aactgctctt ccccctcgtg gcaggccatg    360
150 catgagcctc ggacttttgc tgtatatctt aacaacactg gctacagaac agcctttttt    420
151 ggaaaatacc tcaatgaata taatggcagc tacatccccc ctgggtggcg agaatggctt    480
152 ggattaatca agaattctcg cttctataat tacactgttt gtcgcaatgg catcaaagaa    540
153 aagcatggat ttgattatgc aaaggactac ttcacagact taatcactaa cgagagcatt    600
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154 aattacttca aaatgtctaa gagaatgtat ccccataggc ccgttatgat ggtgatcagc   660
155 cacgctgcgc cccacggccc cgaggactca gccccacagt tttctaaact gtaccccaat   720
156 gcttcccaac acataactcc tagttataac tatgcaccaa atatggataa acactggatt   780
157 atgcagtaca caggaccaat gctgcccatc cacatggaat ttacaaacat tctacagcgc   840
158 aaaaggctcc agactttgat gtcagtggat gattctgtgg agaggctgta taacatgctc   900
159 gtggagacgg gggagctgga gaatacttac atcatttaca ccgccgacca tggttaccat   960
160 attgggcagt ttggactggt caaggggaaa tccatgccat atgactttga tattcgtgtg  1020
161 ccttttttta ttcgtggtcc aagtgtagaa ccaggatcaa tagtcccaca gatcgttctc  1080
162 aacattgact tggcccccac gatcctggat attgctgggc tcgacacacc tcctgatgtg  1140
163 gacggcaagt ctgtcctcaa acttctggac ccagaaaagc caggtaacag gtttcgaaca  1200
164 aacaagaagg ccaaaatttg gcgtgataca ttcctagtgg aaagaggcaa atttctacgt  1260
165 aagaaggaag aatccagcaa gaatatccaa cagtcaaatc acttgcccaa atatgaacgg  1320
166 gtcaaagaac tatgccagca ggccaggtac cagacagcct gtgaacaacc ggggcagaag  1380
167 tggcaatgca ttgaggatac atctggcaag cttcgaattc acaagtgtaa aggacccagt  1440
168 gacctgctca cagtccggca gagcacgcgg aacctctacg ctcgcggctt ccatgacaaa  1500
169 gacaaagagt gcagttgtag ggagtctggt taccgtgcca gcagaagcca agaaagagt   1560
170 caacggcaat tcttgagaaa ccaggggact ccaaagtaca agcccagatt tgtccatact  1620
171 cggcagacac gttccttgtc cgtcgaattt gaaggtgaaa tatatgacat aaatctggaa  1680
172 gaagaagaag aattgcaagt gttgcaacca agaaacattg ctaagcgtca tgatgaaggc  1740
173 cacaaggggc caagagatct ccaggcttcc agtggtggca acaggggcag gatgctggca  1800
174 gatagcagca acgccgtggg cccacctacc actgtccgag tgacacacaa gtgttttatt  1860
175 cttcccaatg actctatcca ttgtgagaga gaactgtacc aatcggccag agcgtggaag  1920
176 gaccataagg catacattga caaagagatt gaagctctgc aagataaaat taagaattta  1980
177 agagaagtga gaggacatct gaagagaagg aagcctgagg aatgtagctg cagtaaacaa  2040
178 agctattaca ataaagagaa aggtgtaaaa aagcaagaga aattaaagag ccatcttcac  2100
179 ccattcaagg aggctgctca ggaagtagat agcaaactgc aactttttcaa ggagaacaac  2160
180 cgtaggagga agaaggagag gaaggagaag agacggcaga ggaaggggga agagtgcagc  2220
181 ctgcctggcc tcacttgctt cacgcatgac aacaaccact ggcagacagc cccgttctgg  2280
182 aacctgggat ctttctgtgc ttgcacgagt tctaacaata acacctactg gtgtttgcgt  2340
183 acagttaatg agacgcataa tttctctttc tgtgagtttg ctactggctt tttggagtat  2400
184 tttgatatga atacagatcc ttatcagctc acaaatacag tgcacacggt agaacgaggc  2460
185 attttgaatc agctacacgt acaactaatg gagctcagaa gctgtcaagg atataagcag  2520
186 tgcaacccaa gacctaagaa tcttgatgtt ggaaataaag atggaggaag ctatgaccta  2580
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189 <210> SEQ ID NO: 3
190 <211> LENGTH: 39
191 <212> TYPE: DNA
192 <213> ORGANISM: Artificial Sequence
194 <220> FEATURE:
195 <223> OTHER INFORMATION: Primer
197 <400> SEQUENCE: 3
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201 <211> LENGTH: 31
202 <212> TYPE: DNA
203 <213> ORGANISM: Artificial Sequence
205 <220> FEATURE:
206 <223> OTHER INFORMATION: Primer
208 <400> SEQUENCE: 4

RAW SEQUENCE LISTING   DATE: 12/28/2004
PATENT APPLICATION: US/10/518,184   TIME: 11:53:52

Input Set : A:\3057US0P.SEQ.txt
Output Set: N:\CRF4\12282004\J518184.raw

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    212 <211> LENGTH: 20
    213 <212> TYPE: DNA
    214 <213> ORGANISM: Artificial Sequence
    216 <220> FEATURE:
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    219 <400> SEQUENCE: 5
    220 gagagaggcg aatggaacga                                        20
    222 <210> SEQ ID NO: 6
    223 <211> LENGTH: 19
    224 <212> TYPE: DNA
    225 <213> ORGANISM: Artificial Sequence
    227 <220> FEATURE:
    228 <223> OTHER INFORMATION: Primer
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    230 cgcaccaggg agctgatct                                         19
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VERIFICATION SUMMARY   DATE: 12/28/2004
PATENT APPLICATION: **US/10/518,184**   TIME: 11:53:53

Input Set : **A:\3057USOP.SEQ.txt**
Output Set: **N:\CRF4\12282004\J518184.raw**

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L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:14 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:17 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:229 M:283 W: Missing Blank Line separator, <400> field identifier